

TableS2. Allele frequencies of 51 SNP loci for 19 *Mytilus* spp. populations

Locus	Allele	ARG30	ARG73	ARG9	IPL	PMD	SAO	BCA	COM	MDP	UBC	PZC	PAR	IRD	LGF	CAM	ORI	NZA	AKAR	KKAT		
BM101A	1	A	0	0	0	0.0714	0.5385	0	0.0147	0	0.0139	0.2414	0.4667	0.2879	0	0.0417	1	1	1	1	0	
BM101A	2	T	1	1	1	0.9286	0.4615	1	0.9853	1	0.9861	0.7586	0.5333	0.7121	1	0.9583	0	0	0	0	1	
BM102A	1	C	0.1167	0.0862	0.1207	0.0345	0.0769	0.0526	0.0735	0.1	0.0139	0.2414	0.3833	0.1818	0.4667	0.3125	0.3393	0.12	0	0	0	
BM102A	2	T	0.8833	0.9138	0.8793	0.9655	0.9231	0.9474	0.9265	0.9	0.9861	0.7586	0.6167	0.8182	0.5333	0.6875	0.6607	0.88	1	1	1	
BM103B	1	A	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	0.2593	
BM103B	2	G	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.7407	
BM105A	1	A	1	1	1	1	1	1	1	1	0.9861	1	1	1	1	0.9815	0.431	0.5714	0.8704	1	1	
BM105A	2	G	0	0	0	0	0.2593	0	0	0	0.0139	0	0	0	0	0.0185	0.569	0.4286	0.1296	0	0	
BM106B	1	A	0	0.0333	0.0333	0	0.3333	0	0.1176	0.1143	0	0	0	0	1	1	1	0.8571	0.2692	0	0.9773	
BM106B	2	G	1	0.9667	0.9667	1	0.6667	1	0.8824	0.8857	1	1	1	1	0	0	0.1429	0.7308	1	0.0227		
BM10B	1	A	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	0.1923	
BM10B	2	C	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.8077		
BM113A	1	A	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	0	
BM113A	2	T	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	
BM118A	1	A	0	0	0	0.1042	0	0	0	0	0	0	0	0	0	0.0577	0.0926	0.2586	0	0	0.0357	
BM118A	2	G	1	1	1	1	0.8958	1	1	1	1	1	1	1	1	0.9423	0.7074	0.7414	1	1	0.9643	
BM11A	1	A	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	0	
BM11A	2	G	0	0	0	0	0	0	0	0	0	0	0	0	0	0.0167	0	0	0	0	1	
BM12A	1	C	0	0	0	0.3519	0	0	0.0147	0	0	0	0	0	0.8167	0.9815	0.9828	1	1	0.9833	0.125	
BM12A	2	T	1	1	1	1	0.6481	1	1	0.9853	1	1	1	1	0.1833	0.0185	0.0172	0	0	0.0167	0.875	
BM12C	1	C	0.9483	1	0.9833	1	0.8654	1	0.9677	0.9839	0.9028	0.9655	1	0.9211	0.2143	0.0926	0.1897	0.7414	0.2963	0	0.6154	
BM12C	2	T	0.0517	0	0.0167	0	0.1346	0	0.0323	0.0161	0.0972	0.0345	0	0.0789	0.7857	0.9074	0.8103	0.2586	0.7037	1	0.3846	
BM147A	1	C	0.9667	0.9833	0.9667	0.8621	0.9259	0.9474	0.8529	0	0.9306	0.6897	0.4667	0.7273	1	0.9	0.931	0.7241	0.3148	0.2	0	
BM147A	2	T	0.0333	0.0167	0.0333	0.1379	0.0741	0.0526	0.1471	0.1	0.0694	0.3103	0.5333	0.2727	0	0	0.1	0.069	0.2759	0.6852	0.8	1
BM151A	1	G	0.9483	0.931	0.9038	0.8966	0.9615	0.8611	0.9853	0.9857	0.9028	0	0	0	1	1	1	1	1	1	0	
BM151A	2	T	0.0517	0.069	0.096	0.1034	0.0385	0.1389	0.0147	0.0143	0.0972	1	1	1	0	0	0	0	1	1	0	
BM16B	1	A	0.89	0.8793	0.8667	0.8621	0.7037	0.8889	0.803	0.8143	0.8611	0.7759	0.7333	0.803	0.8333	0.75	0.7931	0.7931	0.9074	0.95	0.9107	
BM16B	2	G	0.1	0.1207	0.1333	0.1379	0.2963	0.1111	0.197	0.1857	0.1389	0.2241	0.2667	0.197	0.1667	0.25	0.2069	0.2069	0.0926	0.05	0.0893	
BM17B	1	A	1	1	1	0.931	0.48	1	0.9853	1	0.9861	0.7586	0.5333	0.697	0.8667	0.6852	0	0	0.0185	0	0.1	
BM17B	2	G	0	0	0	0.069	0.52	0	0	0.0147	0	0.0139	0.2414	0.4667	0.303	0.1333	0.3148	1	1	0.9815	1	0.9115
BM201B	1	A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.7115	
BM201B	2	C	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	0.2885	
BM201C	1	G	0	0	0	0.38	0	0	0	0	0	0	0	0	0.0167	0	0.0167	0.2759	0.6852	0.8	1	
BM201C	2	T	0	0.0517	0.069	0.096	0.1034	0.0385	0.1389	0.0147	0.0143	0.0972	1	1	1	0	0	0	1	1	0	
BM16B	1	A	0.9	0.8793	0.8667	0.8621	0.7037	0.8889	0.803	0.8143	0.8611	0.7759	0.7333	0.803	0.8333	0.75	0.7931	0.7931	0.9074	0.95	0.9107	
BM203D	1	A	0.85	0.9138	0.9828	0.875	0.3913	0.9722	0.9091	0.8235	0.9697	0.2222	0.0333	0.0806	0.6724	0.4286	0.1296	0.0172	0.1111	0.0517	0	
BM203C	2	G	0.15	0.0862	0.0172	0.125	0.6087	0.0278	0.0909	0.1765	0.0303	0.7778	0.9667	0.9194	0.3276	0.5714	0.8704	0.9828	0.8889	0.9483	1	
BM203D	1	A	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	0	
BM204D	2	T	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	
BM204A	1	C	0	0	0	0	0.2963	0	0	0	0	0	0	0	0.0833	0.0556	0.3571	0.6429	0.2222	0.15	0	
BM204A	2	T	1	1	1	0.7037	1	1	1	1	1	1	1	1	0.9167	0.9444	0.6429	0.3571				